

FIGURE 1 A

210 ACA12 ABE3 290  
AKG7  
MPLPRQNHEPVATSPSSPQPAETHPTTLQGAIIRREPTSSPLYSYTTDGN~~DT~~VTESSDGLWNNNQTQLFLEHSLLTANTTKG  
#43 MPLPRQNHEPVATSPSSP  
#44 PVATSPSSPQPAETHPTT  
#45 QPAETHPTTLQGAIIRREP  
#46 LQGAIIRREPTSSPLYSYT  
#47 TSSPLYSYTTDGN~~DT~~VT  
#48 TDGN~~DT~~VTESSDGLWNNN  
#49 SSDGLWNNNQTQLFLEH  
#50 QTQLFLEHSLLTANTTKG

FIGURE 1B

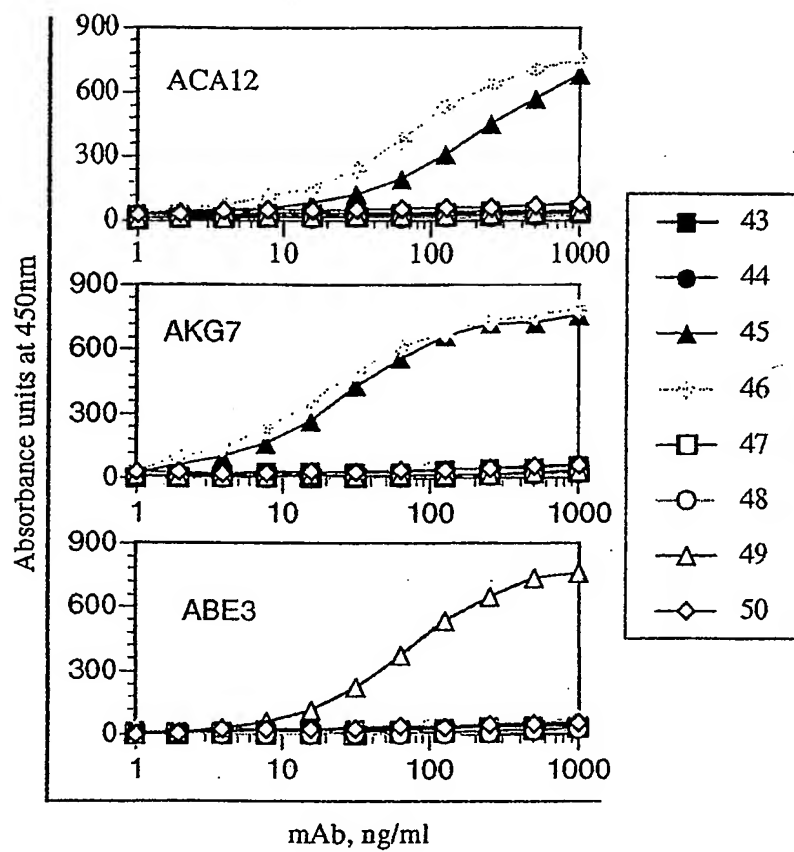


FIGURE 2

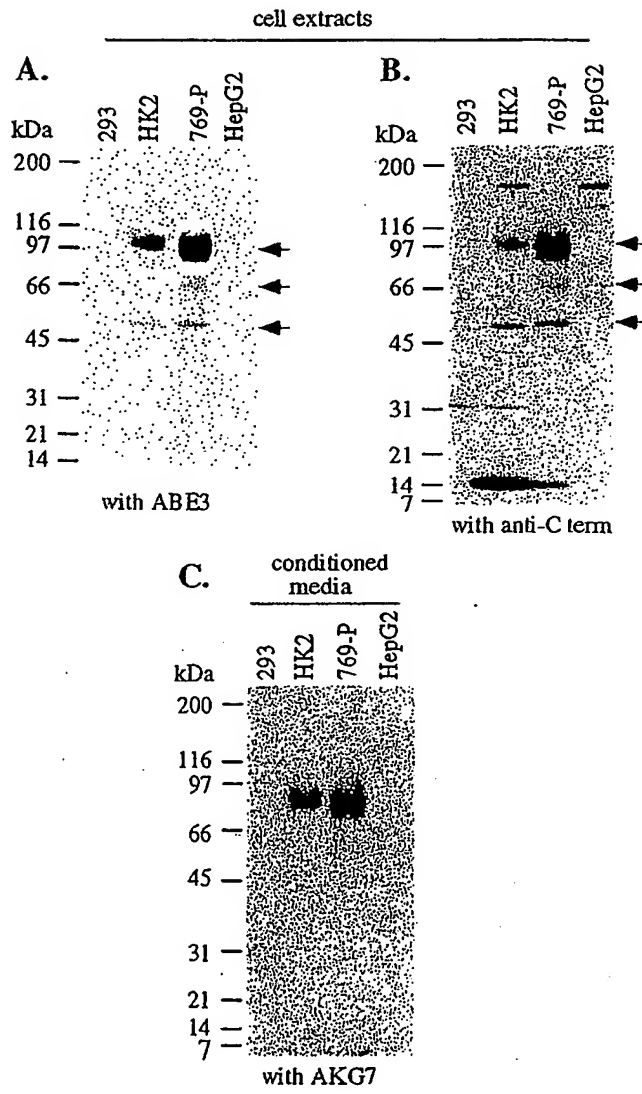


FIGURE 3

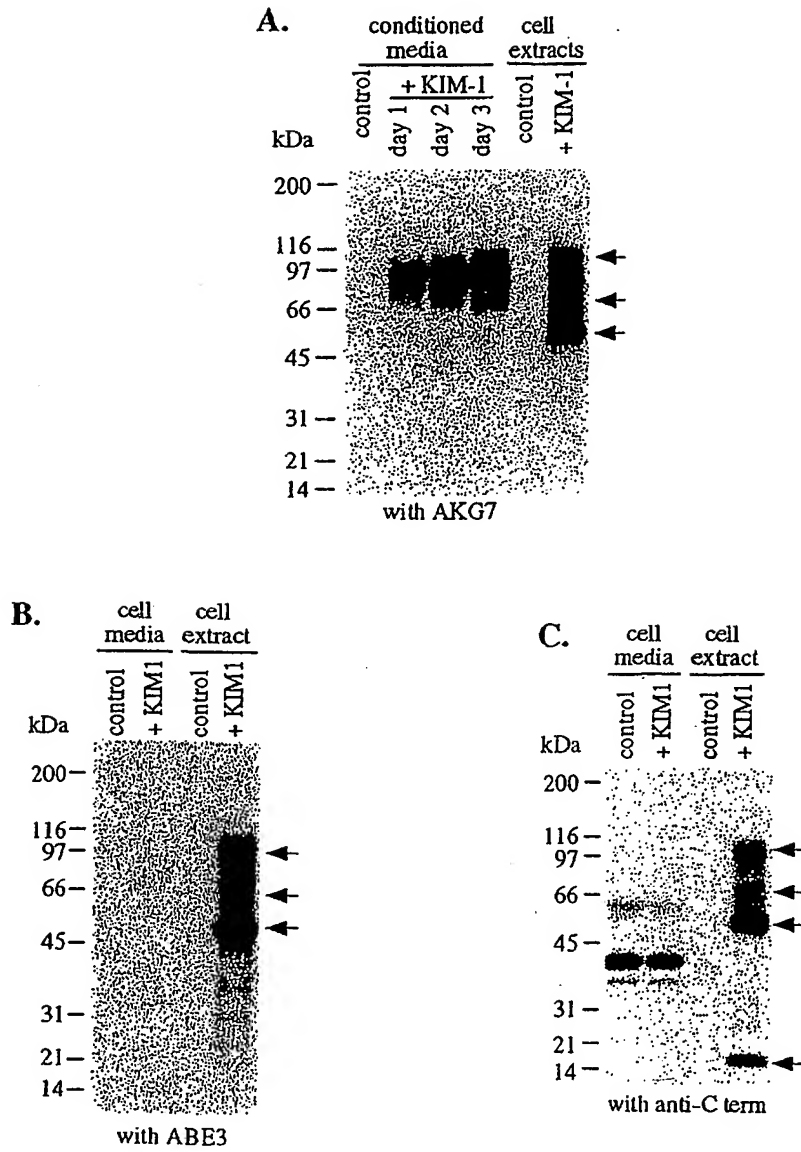


FIGURE 4

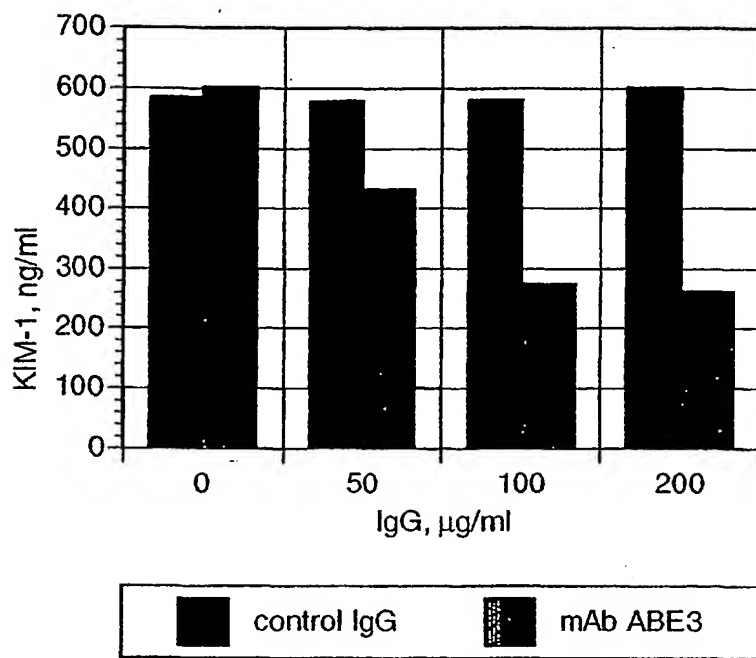
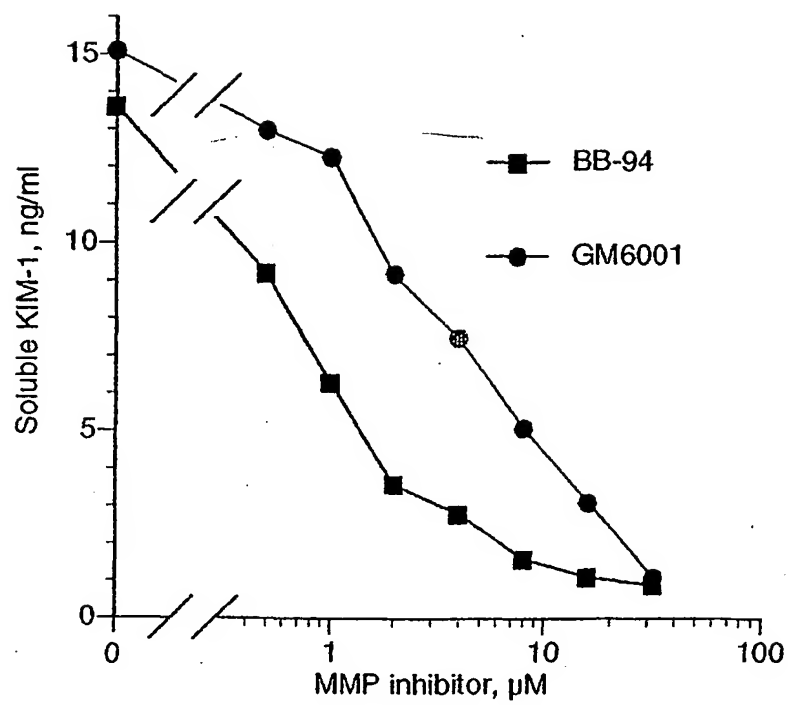


FIGURE 5



**FIGURES 6A and 6B**

A.

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> human KIM-1 (a) 334 aa
> human KIM-1 (b) (HAVcr-1) 359 aa
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MHPOVVL~~SLILHLAD~~SVAGSVKVGGEAGPSVTLPC~~HYSGAVTSMCWN~~RGSCSLFTCQNG 60  
 IVWNG~~THVTYRK~~DTYKLLGDLRRDVS~~LTIENTAVSD~~SGVYCCRVEH~~RGWFND~~MKITV 120  
 SL~~EVPPKVT~~TTTTPIVTVPTVT~~TKSTTVPTTTT~~PTTIVPTTMSIP~~TTTTVPTT~~MTVS 180  
 TTTSVPTTTSIP~~TTTSVPVTTT~~VS~~TFVPMPLPR~~ONHEPVATSPSS~~PQPAETHPTT~~LQGA 240  
 IRREPTSSPLYSY~~TDTG~~DTGV~~TES~~SDGLWNN~~QQLFL~~ELHSL~~LTAT~~TTKGLYAGV~~CLSVL~~ 300  
 VLLALLGVIIAKKYFFKKEVQQL{RPHKSCIQHRE 334  
 VSVSFSSLQIKALQNAVEKEVQ~~AE~~DNIIYIENS~~LYATD~~ 359

**B.**

